



SEQUENCE LISTING

<110> HITACHI, LTD

<120> Method for Testing Nucleic Acid Sequence

<130> WO216-01CW; ASAM.0055

<140> US 10/083,340

<141> 2002-02-27

<150> JP 2001-331853

<151> 2001-10-30

<160> 4

<210> 1

<211> 95

<212> DNA

<213> Artificial Sequence

<220>

<223> Sequence of portion of model gene including mutation.

<400> 1

ctttcttgcg gagattctct tcctctgtgc gccggctct cccaggacag gcacaaacac 60
gcacctcaaa gctgttccgt cccagtagat tacca 95

<210> 2

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> DNA probe for detecting wild type and having an artificial mismatched base at the third base from the 3'end.

<400> 2

aacagctttg aggtgcgtga ta 22

<210> 3

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> DNA probe for detecting mutation type and having an artificial mismatched base at the third base from the 3'end.

<400> 3

aacagctttg aggtgcgtga tt 22

<210> 4

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Sequence portion of SEQ.No.1 which is a target sequence hybridized with DNA probe having base sequence of SEQ.No.2 or

No. 3.

<400> 4
aaacacgcac ctc aaagctg tt 22

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14:01:28

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VERIFICATION SUMMARY REPORT
PATENT APPLICATION

DATE:
TIME:

INPUT SEQ: A:\ASAM55.ans

GENERAL INFORMATION SECTION

3,<110> HITACHI,LTD.
5,<120> Method for Testing Nucleic Acid Sequence
7,<130> WO216-01CW; ASAM.0055
9,<140> US 10/083,340
10,<141> 2002-02-27
12,<150> JP 2001-331853
13,<151> 2001-10-30
15,<160> 4

ERRORED LINES SECTION

STATISTICS SUMMARY

Application Serial Number: US 10/083,340
Alpha or Numeric: Numeric
Application Class:
Application File Date: 2002-02-27
Art Unit:
Software Application:
Total Number of Sequences: 4
Total Nucleotides: 161
Total Amino Acids: 0
Number of Errors: 0
Number of Warnings: 0
Number of Corrections: 0